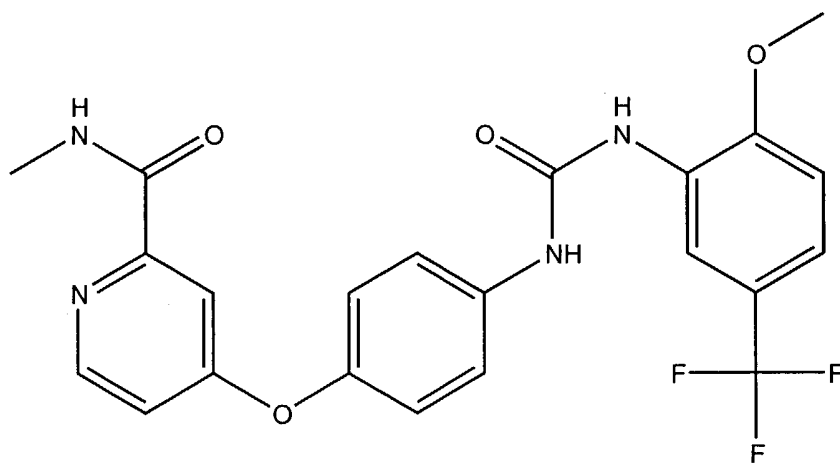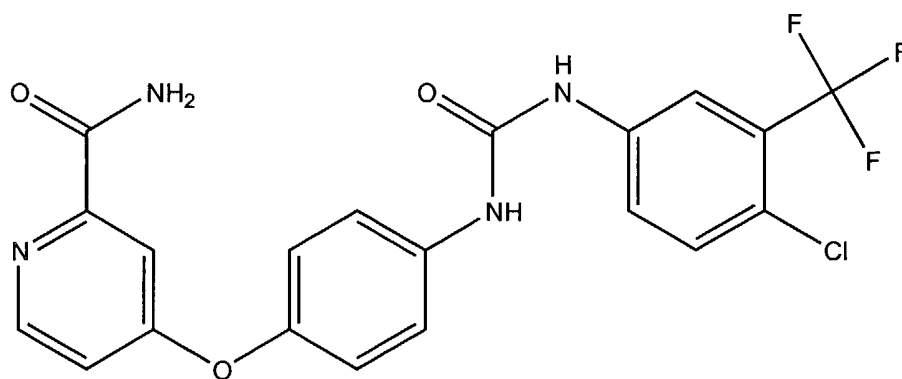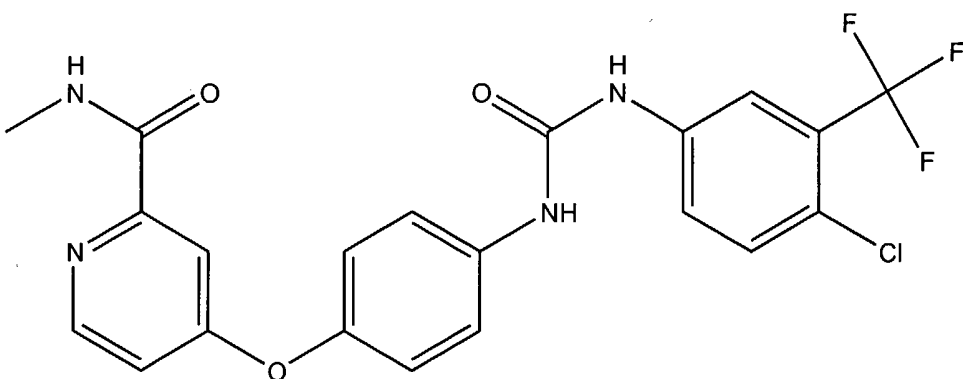N-(5-tert-butyl-2-methoxy phenyl)-N'-(4-(4-methoxy-3(N-methylcarbamoyl)phenoxy)phenyl)urea
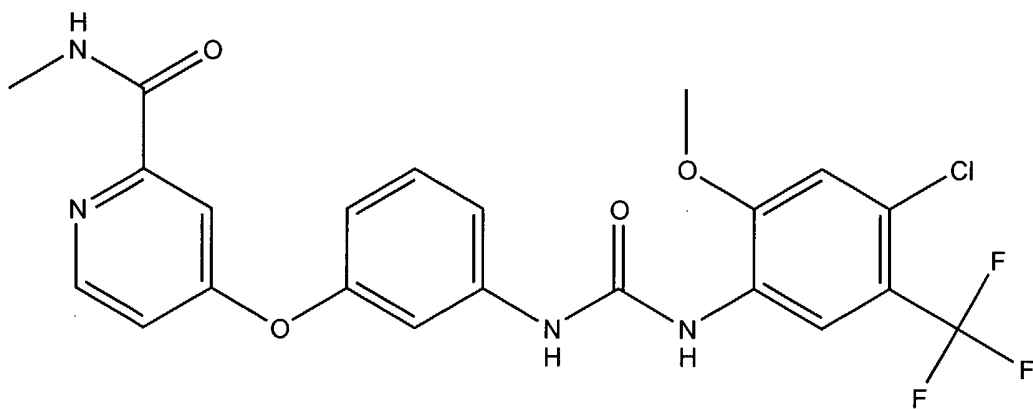

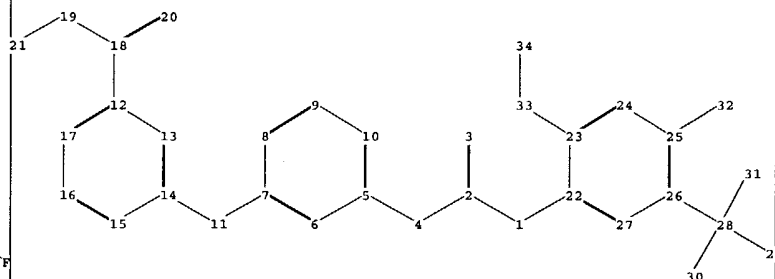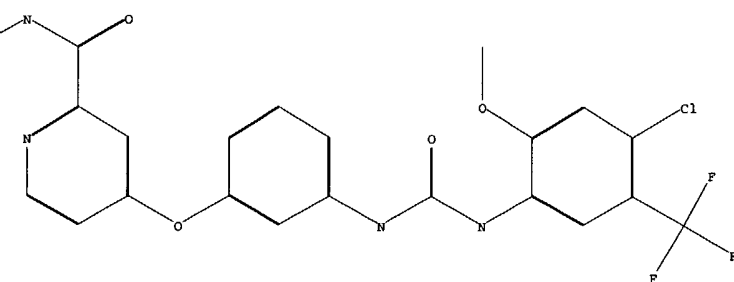N-(2-methoxy -5-(trifluoromethyl)phenyl)-N'-(4-(2-(N-methylcarbamoyl)-4-pyridyloxy)phenyl) urea

N-(4-chloro-3-(trifluoromethyl)phenyl)-N'-(4-(2-carbamoyl-4-pyridyloxy)phenyl) urea

N-(4-chloro-3-(trifluoromethyl)phenyl)-N'-(4-(2(N-methylcarbamoyl)-4-pyridyloxy)phenyl) urea

N-(2-methoxy-4-chloro-5-(trifluoromethyl)phenyl)-N'-(3-(2(N-methylcarbamoyl)-4-pyridyloxy)phenyl) urea

ng nodes :
  5   6   7   8   9   10   12   13   14   15   16   17   22   23   24   25   26   27
ng/chain nodes :
  1   2   3   4   11   18   19   20   21   28   29   30   31   32   33   34
ng/chain bonds :
  1-2   1-22   2-3   2-4   4-5   7-11   11-14   12-18   18-19   18-20   19-21   23-33   25-32   26-28
  28-29   28-30   28-31   33-34
ng bonds :
  5-6   5-10   6-7   7-8   8-9   9-10   12-13   12-17   13-14   14-15   15-16   16-17   22-23   22-27
  23-24   24-25   25-26   26-27
act/norm bonds :
  12-18   25-32   26-28   28-29   28-30   28-31
act bonds :
  1-2   1-22   2-3   2-4   4-5   7-11   11-14   18-19   18-20   19-21   23-33   33-34
rmalized bonds :
  5-6   5-10   6-7   7-8   8-9   9-10   12-13   12-17   13-14   14-15   15-16   16-17   22-23   22-27
  23-24   24-25   25-26   26-27

tch level :
  1:CLASS   2:CLASS   3:CLASS   4:CLASS   5:Atom   6:Atom   7:Atom   8:Atom   9:Atom   10:Atom
  11:CLASS   12:Atom   13:Atom   14:Atom   15:Atom   16:Atom   17:Atom   18:CLASS   19:CLASS
  20:CLASS   21:CLASS   22:Atom   23:Atom   24:Atom   25:Atom   26:Atom   27:Atom   28:CLASS
  29:CLASS   30:CLASS   31:CLASS   32:CLASS   33:CLASS   34:CLASS

ing nodes :
    5   6   7   8   9   10   12   13   14   15   16   17   22   23   24   25   26   27
ing/chain nodes :
    1   2   3   4   11   18   19   20   21   28   29   30   31   32
ing/chain bonds :
    1-2   1-22   2-3   2-4   4-5   8-11   11-14   12-18   18-19   18-20   19-21   24-28   25-32   28-29
    28-30   28-31
ing bonds :
    5-6   5-10   6-7   7-8   8-9   9-10   12-13   12-17   13-14   14-15   15-16   16-17   22-23   22-27
    23-24   24-25   25-26   26-27
xact/norm bonds :
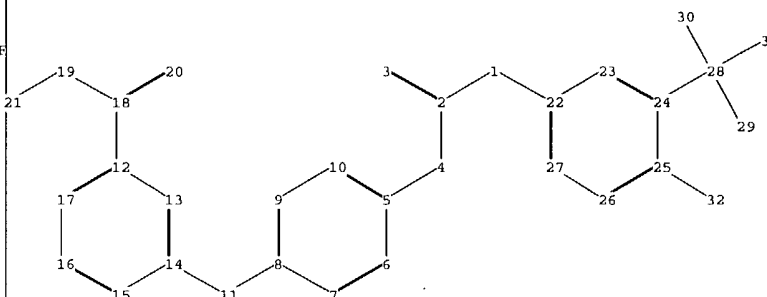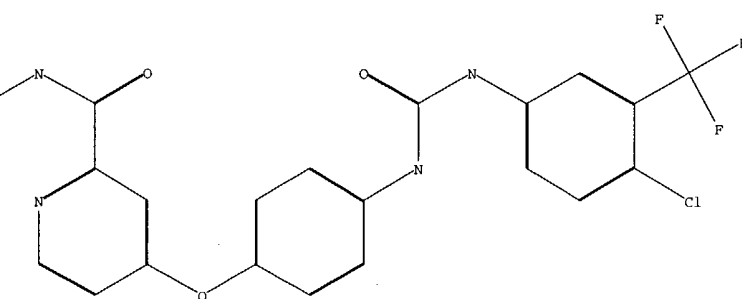   12-18   24-28   25-32   28-29   28-30   28-31
xact bonds :
    1-2   1-22   2-3   2-4   4-5   8-11   11-14   18-19   18-20   19-21
ormalized bonds :
    5-6   5-10   6-7   7-8   8-9   9-10   12-13   12-17   13-14   14-15   15-16   16-17   22-23   22-27
    23-24   24-25   25-26   26-27

atch level :
    1:CLASS   2:CLASS   3:CLASS   4:CLASS   5:Atom   6:Atom   7:Atom   8:Atom   9:Atom   10:Atom
    11:CLASS   12:Atom   13:Atom   14:Atom   15:Atom   16:Atom   17:Atom   18:CLASS   19:CLASS
    20:CLASS   21:CLASS   22:Atom   23:Atom   24:Atom   25:Atom   26:Atom   27:Atom   28:CLASS
    29:CLASS   30:CLASS   31:CLASS   32:CLASS

```
ing nodes :
    5   6   7   8   9   10   12   13   14   15   16   17   21   22   23   24   25   26
ing/chain nodes :
    1   2   3   4   11   18   19   20   27   28   29   30   31
ing/chain bonds :
    1-2   1-21   2-3   2-4   4-5   8-11   11-14   12-18   18-19   18-20   23-27   24-31   27-28   27-29
    27-30
ing bonds :
    5-6   5-10   6-7   7-8   8-9   9-10   12-13   12-17   13-14   14-15   15-16   16-17   21-22   21-26
    22-23   23-24   24-25   25-26
xact/norm bonds :
    12-18   23-27   24-31   27-28   27-29   27-30
xact bonds :
    1-2   1-21   2-3   2-4   4-5   8-11   11-14   18-19   18-20
ormalized bonds :
    5-6   5-10   6-7   7-8   8-9   9-10   12-13   12-17   13-14   14-15   15-16   16-17   21-22   21-26
    22-23   23-24   24-25   25-26

atch level :
    1:CLASS   2:CLASS   3:CLASS   4:CLASS   5:Atom   6:Atom   7:Atom   8:Atom   9:Atom   10:Atom
    11:CLASS   12:Atom   13:Atom   14:Atom   15:Atom   16:Atom   17:Atom   18:CLASS   19:CLASS
    20:CLASS   21:Atom   22:Atom   23:Atom   24:Atom   25:Atom   26:Atom   27:CLASS   28:CLASS
    29:CLASS   30:CLASS   31:CLASS
```
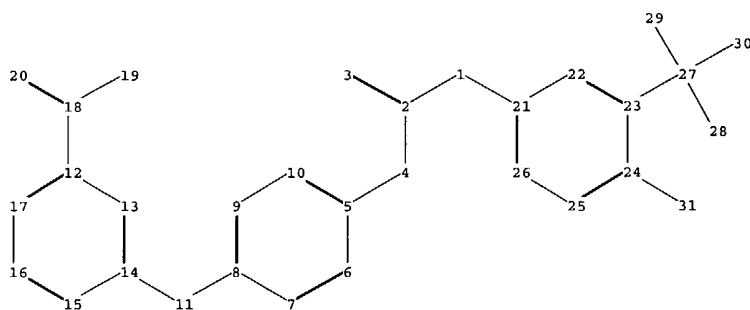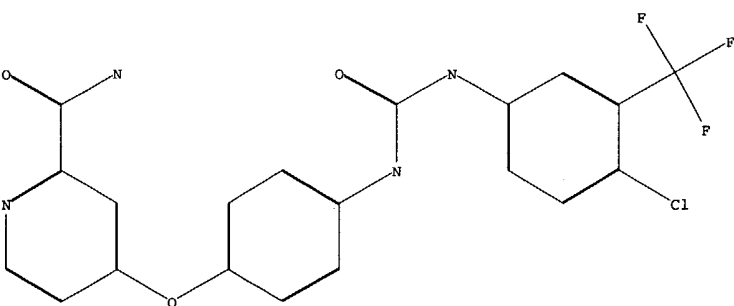
ing nodes :
    5   6   7   8   9   10   12   13   14   15   16   17   22   23   24   25   26   27
ing/chain nodes :
    1   2   3   4   11   18   19   20   21   28   29   30   31   32   33
ing/chain bonds :
    1-2   1-22   2-3   2-4   4-5   8-11   11-14   12-18   18-19   18-20   19-21   23-32   26-28   28-29
    28-30   28-31   32-33
ing bonds :
    5-6   5-10   6-7   7-8   8-9   9-10   12-13   12-17   13-14   14-15   15-16   16-17   22-23   22-27
    23-24   24-25   25-26   26-27
xact/norm bonds :
    12-18   26-28   28-29   28-30   28-31
xact bonds :
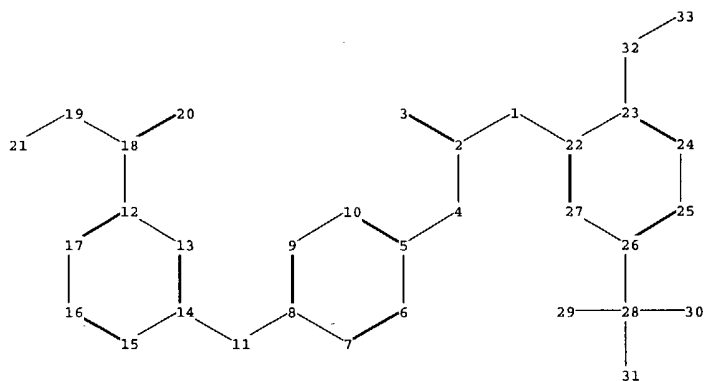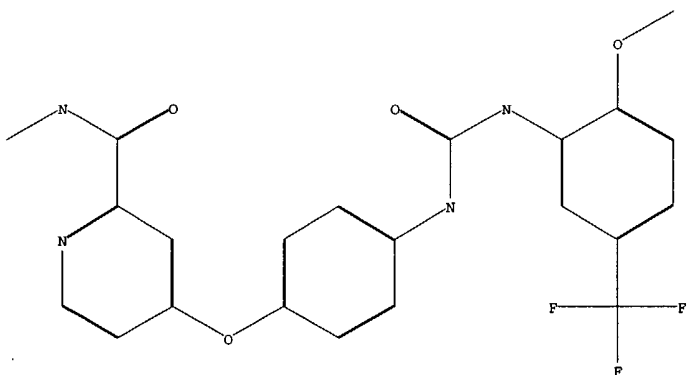   1-2   1-22   2-3   2-4   4-5   8-11   11-14   18-19   18-20   19-21   23-32   32-33
ormalized bonds :
    5-6   5-10   6-7   7-8   8-9   9-10   12-13   12-17   13-14   14-15   15-16   16-17   22-23   22-27
    23-24   24-25   25-26   26-27

atch level :
    1:CLASS   2:CLASS   3:CLASS   4:CLASS   5:Atom   6:Atom   7:Atom   8:Atom   9:Atom   10:Atom
    11:CLASS   12:Atom   13:Atom   14:Atom   15:Atom   16:Atom   17:Atom   18:CLASS   19:CLASS
    20:CLASS   21:CLASS   22:Atom   23:Atom   24:Atom   25:Atom   26:Atom   27:Atom   28:CLASS
    29:CLASS   30:CLASS   31:CLASS   32:CLASS   33:CLASS

ng nodes :
    5   6   7   8   9   10   12   13   14   15   16   17   24   25   26   27   28   29
ng/chain nodes :
    1   2   3   4   11   18   19   20   21   22   23   30   31   32   33   34   35
ng/chain bonds :
    1-2   1-24   2-3   2-4   4-5   8-11   11-12   14-18   15-22   18-19   18-20   19-21   22-23   25-30
    28-32   30-31   32-33   32-35   32-34
ng bonds :
    5-6   5-10   6-7   7-8   8-9   9-10   12-13   12-17   13-14   14-15   15-16   16-17   24-25   24-29
    25-26   26-27   27-28   28-29
act/norm bonds :
    14-18   28-32   32-33   32-35   32-34
act bonds :
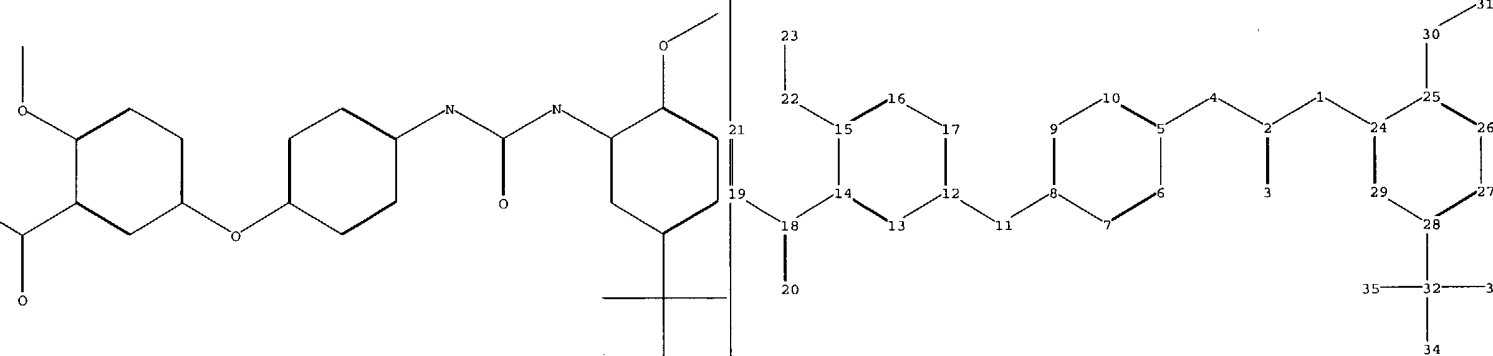   1-2   1-24   2-3   2-4   4-5   8-11   11-12   15-22   18-19   18-20   19-21   22-23   25-30   30-31
rmalized bonds :
    5-6   5-10   6-7   7-8   8-9   9-10   12-13   12-17   13-14   14-15   15-16   16-17   24-25   24-29
    25-26   26-27   27-28   28-29

tch level :
    1:CLASS   2:CLASS   3:CLASS   4:CLASS   5:Atom   6:Atom   7:Atom   8:Atom   9:Atom   10:Atom
    11:CLASS   12:Atom   13:Atom   14:Atom   15:Atom   16:Atom   17:Atom   18:CLASS   19:CLASS
    20:CLASS   21:CLASS   22:CLASS   23:CLASS   24:Atom   25:Atom   26:Atom   27:Atom   28:Atom
    29:Atom   30:CLASS   31:CLASS   32:CLASS   33:CLASS   34:CLASS   35:CLASS

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PASSWORD:
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10042203

8/16/04

FILE 'HOME' ENTERED AT 18:25:12 ON 16 AUG 2004

=> file registry
COST IN U.S. DOLLARS   SINCE FILE   TOTAL
                                                    ENTRY   SESSION
FULL ESTIMATED COST   0.21   0.21

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DICTIONARY FILE UPDATES: 15 AUG 2004   HIGHEST RN 727358-71-6

TSCA INFORMATION NOW CURRENT THROUGH MAY 21, 2004

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=>
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L1   STRUCTURE UPLOADED

=> d l1
L1 HAS NO ANSWERS
L1   STR

* STRUCTURE DIAGRAM TOO LARGE, FOR DISPLAY - AVAILABLE VIA OFFLINE PRINT *

Structure attributes must be viewed using STN Express query preparation.

=> s l1
SAMPLE SEARCH INITIATED 18:25:41 FILE 'REGISTRY'
SAMPLE SCREEN SEARCH COMPLETED -   142 TO ITERATE

100.0% PROCESSED   142 ITERATIONS   0 ANSWERS
SEARCH TIME: 00.00.01

FULL FILE PROJECTIONS: ONLINE   **COMPLETE**
                        BATCH   **COMPLETE**
PROJECTED ITERATIONS: 2126 TO   3554
PROJECTED ANSWERS: 0 TO   0

L2   0 SEA SSS SAM L1

=> s l1 ful

10042203

8/16/04

FULL SEARCH INITIATED 18:25:45 FILE 'REGISTRY'
FULL SCREEN SEARCH COMPLETED -   2905 TO ITERATE

100.0% PROCESSED   2905 ITERATIONS   0 ANSWERS
SEARCH TIME: 00.00.01

L3   0 SEA SSS FUL L1

=> file registry
COST IN U.S. DOLLARS   SINCE FILE   TOTAL
                                              ENTRY   SESSION
FULL ESTIMATED COST   156.26   156.47

FILE 'REGISTRY' ENTERED AT 18:27:01 ON 16 AUG 2004
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DICTIONARY FILE UPDATES: 15 AUG 2004   HIGHEST RN 727358-71-6

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=>
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L4   STRUCTURE UPLOADED

=> d 14
L4 HAS NO ANSWERS
L4   STR

* STRUCTURE DIAGRAM TOO LARGE FOR DISPLAY -   AVAILABLE VIA OFFLINE PRINT *

Structure attributes must be viewed using STN Express query preparation.

=> s 14
SAMPLE SEARCH INITIATED 18:27:23 FILE 'REGISTRY'
SAMPLE SCREEN SEARCH COMPLETED -   48 TO ITERATE

100.0% PROCESSED   48 ITERATIONS   0 ANSWERS
SEARCH TIME: 00.00.01

FULL FILE PROJECTIONS: ONLINE   **COMPLETE**
                         BATCH   **COMPLETE**

10042203

```
PROJECTED ITERATIONS:              545 TO      1375
PROJECTED ANSWERS:                   0 TO         0

L5              0 SEA SSS SAM L4

=> s l4 ful
FULL SEARCH INITIATED 18:27:28 FILE 'REGISTRY'
FULL SCREEN SEARCH COMPLETED -    803 TO ITERATE

100.0% PROCESSED     803 ITERATIONS                          0 ANSWERS
SEARCH TIME: 00.00.01

L6              0 SEA SSS FUL L4

=> file registry
COST IN U.S. DOLLARS                          SINCE FILE       TOTAL
                                                ENTRY         SESSION
FULL ESTIMATED COST                            157.10         313.57

FILE 'REGISTRY' ENTERED AT 18:29:56 ON 16 AUG 2004
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Experimental and calculated property data are now available. For more
information enter HELP PROP at an arrow prompt in the file or refer
to the file summary sheet on the web at:
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=>
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L7      STRUCTURE UPLOADED

=> d l7
L7 HAS NO ANSWERS
L7                STR

* STRUCTURE DIAGRAM TOO LARGE FOR DISPLAY -  AVAILABLE VIA OFFLINE PRINT *

Structure attributes must be viewed using STN Express query preparation.


=> s l7 ful
FULL SEARCH INITIATED 18:30:26 FILE 'REGISTRY'
FULL SCREEN SEARCH COMPLETED -    3635 TO ITERATE
```

10042203

```
100.0% PROCESSED    3635 ITERATIONS                          0 ANSWERS
SEARCH TIME: 00.00.01


L8              0 SEA SSS FUL L7

=> file registry
COST IN U.S. DOLLARS                           SINCE FILE      TOTAL
                                                 ENTRY        SESSION
FULL ESTIMATED COST                              157.10       470.67

FILE 'REGISTRY' ENTERED AT 18:32:50 ON 16 AUG 2004
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DICTIONARY FILE UPDATES:   15 AUG 2004   HIGHEST RN 727358-71-6

TSCA INFORMATION NOW CURRENT THROUGH MAY 21, 2004

   Please note that search-term pricing does apply when
   conducting SmartSELECT searches.

Crossover limits have been increased.  See HELP CROSSOVER for details.

Experimental and calculated property data are now available. For more
information enter HELP PROP at an arrow prompt in the file or refer
to the file summary sheet on the web at:
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=>
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L9        STRUCTURE UPLOADED

=> d 19
L9 HAS NO ANSWERS
L9                STR

* STRUCTURE DIAGRAM TOO LARGE FOR DISPLAY -  AVAILABLE VIA OFFLINE PRINT *

Structure attributes must be viewed using STN Express query preparation.


=> s 19
SAMPLE SEARCH INITIATED 18:33:12 FILE 'REGISTRY'
SAMPLE SCREEN SEARCH COMPLETED -     167 TO ITERATE

100.0% PROCESSED     167 ITERATIONS                          0 ANSWERS
SEARCH TIME: 00.00.01

FULL FILE PROJECTIONS:   ONLINE   **COMPLETE**
                         BATCH    **COMPLETE**
PROJECTED ITERATIONS:         2565 TO       4115
PROJECTED ANSWERS:               0 TO          0
```

L10   0 SEA SSS SAM L9

=> s l9 ful
FULL SEARCH INITIATED 18:33:17 FILE 'REGISTRY'
FULL SCREEN SEARCH COMPLETED -   3635 TO ITERATE

100.0% PROCESSED   3635 ITERATIONS   0 ANSWERS
SEARCH TIME: 00.00.01

L11   0 SEA SSS FUL L9

=> file registry
COST IN U.S. DOLLARS   SINCE FILE   TOTAL
                                                  ENTRY   SESSION
FULL ESTIMATED COST   157.10   627.77

FILE 'REGISTRY' ENTERED AT 18:35:36 ON 16 AUG 2004
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DICTIONARY FILE UPDATES: 15 AUG 2004   HIGHEST RN 727358-71-6

TSCA INFORMATION NOW CURRENT THROUGH MAY 21, 2004

   Please note that search-term pricing does apply when
   conducting SmartSELECT searches.

Crossover limits have been increased. See HELP CROSSOVER for details.

Experimental and calculated property data are now available. For more
information enter HELP PROP at an arrow prompt in the file or refer
to the file summary sheet on the web at:
http://www.cas.org/ONLINE/DBSS/registryss.html

=>
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L12   STRUCTURE UPLOADED

=> d l12
L12 HAS NO ANSWERS
L12   STR

* STRUCTURE DIAGRAM TOO LARGE FOR DISPLAY - AVAILABLE VIA OFFLINE PRINT *

Structure attributes must be viewed using STN Express query preparation.

=> s l12   .
SAMPLE SEARCH INITIATED 18:36:07 FILE 'REGISTRY'
SAMPLE SCREEN SEARCH COMPLETED -   23 TO ITERATE

100.0% PROCESSED   23 ITERATIONS   0 ANSWERS

10042203

8/16/04

SEARCH TIME: 00.00.01

FULL FILE PROJECTIONS: ONLINE   **COMPLETE**
                         BATCH   **COMPLETE**
PROJECTED ITERATIONS: 173 TO   747
PROJECTED ANSWERS: 0 TO   0

L13   0 SEA SSS SAM L12

=> s l12 ful
FULL SEARCH INITIATED 18:36:12 FILE 'REGISTRY'
FULL SCREEN SEARCH COMPLETED -   354 TO ITERATE

100.0% PROCESSED   354 ITERATIONS   0 ANSWERS
SEARCH TIME: 00.00.01

L14   0 SEA SSS FUL L12

=>